

OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/729,838

DATE: 09/07/2001
TIME: 16:40:43

Input Set : A:\010342-012-999.txt
Output Set: N:\CRF3\09072001\I729838.raw

5 <110> APPLICANT: Lewis, Hal A
9 <120> TITLE OF INVENTION: Crystals and Structures of Luxs
13 <130> FILE REFERENCE: 010342-0012-999
C--> 17 <140> CURRENT APPLICATION NUMBER: US/09/729,838
C--> 17 <141> CURRENT FILING DATE: 2000-12-04
17 <150> PRIOR APPLICATION NUMBER: 60/237,933
19 <151> PRIOR FILING DATE: 2000-10-03
23 <160> NUMBER OF SEQ ID NOS: 13
27 <170> SOFTWARE: PatentIn version 3.1
31 <210> SEQ ID NO: 1
33 <211> LENGTH: 152
35 <212> TYPE: PRT
37 <213> ORGANISM: H. Pylori
41 <400> SEQUENCE: 1
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47 Ala Pro Tyr Val Arg Ile Ala Asp Arg Lys Lys Gly Val Asn Gly Asp
48 20 25 30
51 Leu Ile Val Lys Tyr Asp Val Arg Phe Lys Gln Pro Asn Arg Asp His
52 35 40 45
55 Met Asp Met Pro Ser Leu His Ser Leu Glu His Leu Val Ala Glu Ile
56 50 55 60
59 Ile Arg Asn His Ala Asn Tyr Val Val Asp Trp Ser Pro Met Gly Cys
60 65 70 75 80
63 Gln Thr Gly Phe Tyr Leu Thr Val Leu Asn His Asp Asn Tyr Thr Glu
64 85 90 95
67 Ile Leu Glu Val Leu Glu Lys Thr Met Gln Asp Val Leu Lys Ala Lys
68 100 105 110
71 Glu Val Pro Ala Ser Asn Glu Lys Gln Cys Gly Trp Ala Ala Asn His
72 115 120 125
75 Thr Leu Glu Gly Ala Gln Asn Leu Ala Arg Ala Phe Leu Asp Lys Arg
76 130 135 140
79 Ala Glu Trp Ser Glu Val Gly Val
80 145 150
83 <210> SEQ ID NO: 2
85 <211> LENGTH: 167
87 <212> TYPE: PRT
89 <213> ORGANISM: H. Influenzae
93 <400> SEQUENCE: 2
95 Met Pro Leu Leu Asp Ser Phe Lys Val Asp His Thr Lys Met Asn Ala
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99 Pro Ala Val Arg Ile Ala Lys Thr Met Leu Thr Pro Lys Gly Asp Asn
100 20 25 30
103 Ile Thr Val Phe Asp Leu Arg Phe Cys Ile Pro Asn Lys Glu Ile Leu
104 35 40 45
107 Ser Pro Lys Gly Ile His Thr Leu Glu His Leu Phe Ala Gly Phe Met
108 50 55 60

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111 Arg Asp His Leu Asn Gly Asp Ser Ile Glu Ile Ile Asp Ile Ser Pro
112 65 70 75 80
115 Met Gly Cys Arg Thr Gly Phe Tyr Met Ser Leu Ile Gly Thr Pro Asn
116 85 90 95
119 Glu Gln Lys Val Ser Glu Ala Trp Leu Ala Ser Met Gln Asp Val Leu
120 100 105 110
123 Gly Val Gln Asp Gln Ala Ser Ile Pro Glu Leu Asn Ile Tyr Gln Cys
124 115 120 125
127 Gly Ser Tyr Thr Glu His Ser Leu Glu Asp Ala His Glu Ile Ala Lys
128 130 135 140
131 Asn Val Ile Ala Arg Gly Ile Gly Val Asn Lys Asn Glu Asp Leu Ser
132 145 150 155 160
135 Leu Asp Asn Ser Leu Leu Lys
136 165
139 <210> SEQ ID NO: 3
141 <211> LENGTH: 158
143 <212> TYPE: PRT
145 <213> ORGANISM: D. Radiodurans
149 <400> SEQUENCE: 3
151 Met Pro Asp Met Ala Asn Val Glu Ser Phe Asp Leu Asp His Thr Lys
152 1 5 10 15
155 Val Lys Ala Pro Tyr Val Arg Leu Ala Gly Val Lys Thr Thr Pro Lys
156 20 25 30
159 Gly Asp Gln Ile Ser Lys Tyr Asp Leu Arg Phe Leu Gln Pro Asn Gln
160 35 40 45
163 Gly Ala Ile Asp Pro Ala Ala Ile His Thr Leu Glu His Leu Leu Ala
164 50 55 60
167 Gly Tyr Met Arg Asp His Leu Glu Gly Val Val Asp Val Ser Pro Met
168 65 70 75 80
171 Gly Cys Arg Thr Gly Met Tyr Met Ala Val Ile Gly Glu Pro Asp Glu
172 85 90 95
175 Gln Gly Val Met Lys Ala Phe Glu Ala Ala Leu Lys Asp Thr Ala Gly
176 100 105 110
179 His Asp Gln Pro Ile Pro Gly Val Ser Glu Leu Glu Cys Gly Asn Tyr
180 115 120 125
183 Arg Asp His Asp Leu Ala Ala Ala Arg Gln His Ala Arg Asp Val Leu
184 130 135 140
187 Asp Gln Gly Leu Lys Val Gln Glu Thr Ile Leu Leu Glu Arg
188 145 150 155
191 <210> SEQ ID NO: 4
193 <211> LENGTH: 164
195 <212> TYPE: PRT
197 <213> ORGANISM: C. Jejuni
201 <400> SEQUENCE: 4
203 Met Pro Leu Leu Asp Ser Phe Lys Val Asp His Thr Lys Met Pro Ala
204 1 5 10 15
207 Pro Ala Val Arg Leu Ala Lys Val Met Lys Thr Pro Lys Gly Asp Asp
208 20 25 30
211 Ile Ser Val Phe Asp Leu Arg Phe Cys Ile Pro Asn Lys Asp Ile Met

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212 35 40 45
 215 Ser Glu Lys Gly Thr His Thr Leu Glu His Leu Phe Ala Gly Phe Met
 216 50 55 60
 219 Arg Asp His Leu Asn Ser Asn Ser Val Glu Ile Ile Asp Ile Ser Pro
 220 65 70 75 80
 223 Met Gly Cys Arg Thr Gly Phe Tyr Met Ser Leu Ile Gly Thr Pro Asp
 224 85 90 95
 227 Glu Lys Ser Ile Ala Lys Ala Trp Glu Ala Ala Met Lys Asp Val Leu
 228 100 105 110
 231 Ser Val Ser Asp Gln Ser Lys Ile Pro Glu Leu Asn Ile Tyr Gln Cys
 232 115 120 125
 235 Gly Thr Cys Ala Met His Ser Leu Asp Glu Ala Lys Gln Ile Ala Gln
 236 130 135 140
 239 Lys Val Leu Asn Leu Gly Ile Ser Ile Ile Asn Asn Lys Arg Leu Lys
 240 145 150 155 160
 243 Leu Glu Asn Ala
 247 <210> SEQ ID NO: 5
 249 <211> LENGTH: 157
 251 <212> TYPE: PRT
 253 <213> ORGANISM: B. Burgdorferi
 257 <400> SEQUENCE: 5
 259 Met Lys Ile Thr Ser Phe Thr Ile Asp His Thr Lys Leu Asn Pro
 260 1 5 10 15
 263 Gly Ile Tyr Val Ser Arg Lys Asp Thr Phe Glu Asn Val Ile Phe Thr
 264 20 25 30
 267 Thr Ile Asp Ile Arg Ile Lys Ala Pro Asn Ile Glu Pro Ile Ile Glu
 268 35 40 45
 271 Asn Ala Ala Ile His Thr Ile Glu His Ile Gly Ala Thr Leu Leu Arg
 272 50 55 60
 275 Asn Asn Glu Val Trp Thr Glu Lys Ile Val Tyr Phe Gly Pro Met Gly
 276 65 70 75 80
 279 Cys Arg Thr Gly Phe Tyr Leu Ile Ile Phe Gly Asp Tyr Glu Ser Lys
 280 85 90 95
 283 Asp Leu Val Asp Leu Val Ser Trp Leu Phe Ser Glu Ile Val Asn Phe
 284 100 105 110
 287 Ser Glu Pro Ile Pro Gly Ala Ser Asp Lys Glu Cys Gly Asn Tyr Lys
 288 115 120 125
 291 Glu His Asn Leu Asp Met Ala Lys Tyr Glu Ser Ser Lys Tyr Leu Gln
 292 130 135 140
 295 Ile Leu Asn Asn Ile Lys Glu Glu Asn Leu Lys Tyr Pro
 296 145 150 155
 299 <210> SEQ ID NO: 6
 301 <211> LENGTH: 151
 303 <212> TYPE: PRT
 305 <213> ORGANISM: C. Perfringens
 309 <400> SEQUENCE: 6
 311 Met Val Lys Val Glu Ser Phe Glu Leu Asp His Thr Lys Val Lys Ala
 312 1 5 10 15
 315 Pro Tyr Val Arg Lys Ala Gly Ile Lys Ile Gly Pro Lys Gly Asp Ile

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316 20 25 30
 319 Val Ser Lys Phe Asp Leu Arg Phe Val Gln Pro Asn Lys Glu Leu Leu
 320 35 40 45
 323 Ser Asp Lys Gly Met His Thr Leu Glu His Phe Leu Ala Gly Phe Met
 324 50 55 60
 327 Arg Glu Lys Leu Asp Asp Val Ile Asp Ile Ser Pro Met Gly Cys Lys
 328 65 70 75 80
 331 Thr Gly Phe Tyr Leu Thr Ser Phe Gly Asp Ile Asp Val Lys Asp Ile
 332 85 90 95
 335 Ile Glu Ala Leu Glu Tyr Ser Leu Ser Lys Val Leu Glu Gln Glu Glu
 336 100 105 110
 339 Ile Pro Ala Ala Asn Glu Leu Gln Cys Gly Ser Ala Lys Leu His Ser
 340 115 120 125
 343 Leu Glu Ala Lys Ser His Ala Lys Gln Val Leu Glu Asn Gly Ile
 344 130 135 140
 347 Ser Asp Lys Phe Tyr Val Glu
 348 145 150
 351 <210> SEQ ID NO: 7
 353 <211> LENGTH: 168
 355 <212> TYPE: PRT
 357 <213> ORGANISM: N. Meningitidis
 361 <400> SEQUENCE: 7
 363 Met Pro Leu Leu Asp Ser Phe Lys Val Asp His Thr Arg Met His Ala
 364 1 5 10 15
 367 Pro Ala Val Arg Val Ala Lys Thr Met Thr Thr Pro Lys Gly Asp Thr
 368 20 25 30
 371 Ile Thr Val Phe Asp Leu Arg Phe Cys Val Pro Asn Lys Glu Ile Leu
 372 35 40 45
 375 Pro Glu Lys Gly Ile His Thr Leu Glu His Leu Phe Ala Gly Phe Met
 376 50 55 60
 379 Arg Asp His Leu Asn Gly Asn Gly Val Glu Ile Ile Asp Ile Ser Pro
 380 65 70 75 80
 383 Met Gly Cys Arg Thr Gly Phe Tyr Met Ser Leu Ile Gly Thr Pro Ser
 384 85 90 95
 387 Glu Gln Gln Val Ala Asp Ala Trp Leu Ala Ser Met Gln Asp Val Leu
 388 100 105 110
 391 Asn Val Lys Asp Gln Ser Lys Ile Pro Glu Leu Asn Glu Tyr Gln Cys
 392 115 120 125
 395 Gly Thr Tyr Gln Met His Ser Leu Ala Glu Ala Gln Gln Ile Ala Gln
 396 130 135 140
 399 Asn Val Leu Ala Arg Lys Val Ala Val Asn Lys Asn Glu Leu Thr
 400 145 150 155 160
 403 Leu Asp Glu Gly Leu Leu Asn Ala
 404 165
 407 <210> SEQ ID NO: 8
 409 <211> LENGTH: 171
 411 <212> TYPE: PRT
 413 <213> ORGANISM: S. Typhimurium
 417 <400> SEQUENCE: 8

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419 Met Pro Leu Leu Asp Ser Phe Ala Val Asp His Thr Arg Met Gln Ala
 420 1 5 10 15
 423 Pro Ala Val Arg Val Ala Lys Thr Met Asn Thr Pro His Gly Asp Ala
 424 20 25 30
 427 Ile Thr Val Phe Asp Leu Arg Phe Cys Ile Pro Asn Lys Glu Val Met
 428 35 40 45
 431 Pro Glu Lys Gly Ile His Thr Leu Glu His Leu Phe Ala Gly Phe Met
 432 50 55 60
 435 Arg Asp His Leu Asn Gly Asn Gly Val Glu Ile Ile Asp Ile Ser Pro
 436 65 70 75 80
 439 Met Gly Cys Arg Thr Gly Phe Tyr Met Ser Leu Ile Gly Thr Pro Asp
 440 85 90 95
 443 Glu Gln Arg Val Ala Asp Ala Trp Lys Ala Ala Met Ala Asp Val Leu
 444 100 105 110
 447 Lys Val Gln Asp Gln Asn Gln Ile Pro Glu Leu Asn Val Tyr Gln Cys
 448 115 120 125
 451 Gly Thr Tyr Gln Met His Ser Leu Ser Glu Ala Gln Asp Ile Ala Arg
 452 130 135 140
 455 His Ile Leu Glu Arg Asp Val Arg Val Asn Ser Asn Lys Glu Leu Ala
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 459 Leu Pro Lys Glu Lys Leu Gln Glu Leu His Ile
 460 165 170
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 467 <212> TYPE: PRT
 469 <213> ORGANISM: V. Harveyi
 473 <400> SEQUENCE: 9
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 479 Pro Ala Val Arg Val Ala Lys Thr Met Gln Thr Pro Lys Gly Asp Thr
 480 20 25 30
 483 Ile Thr Val Phe Asp Leu Arg Phe Thr Ala Pro Asn Lys Asp Ile Leu
 484 35 40 45
 487 Ser Glu Lys Gly Ile His Thr Leu Glu His Leu Tyr Ala Gly Phe Met
 488 50 55 60
 491 Arg Asn His Leu Asn Gly Asp Ser Val Glu Ile Ile Asp Ile Ser Pro
 492 65 70 75 80
 495 Met Gly Cys Arg Thr Gly Phe Tyr Met Ser Leu Ile Gly Thr Pro Ser
 496 85 90 95
 499 Glu Gln Gln Val Ala Asp Ala Trp Ile Ala Ala Met Glu Asp Val Leu
 500 100 105 110
 503 Lys Val Glu Asn Gln Asn Lys Ile Pro Glu Leu Asn Glu Tyr Gln Cys
 504 115 120 125
 507 Gly Thr Ala Ala Met His Ser Leu Asp Glu Ala Lys Gln Ile Ala Lys
 508 130 135 140
 511 Asn Ile Leu Glu Val Gly Val Ala Val Asn Lys Asn Asp Glu Leu Ala
 512 145 150 155 160
 515 Leu Pro Glu Ser Met Leu Arg Glu Leu Arg Ile Asp
 516 165 170

VERIFICATION SUMMARY

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Input Set : A:\010342-012-999.txt

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L:17 M:270 C: Current Application Number differs, Replaced Current Application No

L:17 M:271 C: Current Filing Date differs, Replaced Current Filing Date